In accordance with 37 C.F.R. §1.121, Applicant has provided (1) accurate instructions to amend the paragraphs, (2) replacement paragraphs in clean form herein, and (3) another version of the replacement paragraphs marked up to show all the changes relative to the previous version, which appears on an attached page.

I hereby state that the content of the paper and computer readable copies of the Sequence Listing submitted in accordance with 37 C.F.R. § 1.821(c) and (e), respectively, are the same.

I hereby state that the content of the paper and computer readable copies of the Sequence Listing, submitted herewith in accordance with 37 C.F.R. § 1.82(f), does not include new matter.

It is believed that no fee is required in connection with this communication. However, if any fee is required in connection with this communication, the Commissioner is hereby authorized to charge payment of any such fee to Deposit Account No. 02-4377. Two copies of this paper are enclosed.

Respectfully submitted,

Louis S. Sorell
PTO Reg. No. 32,439

Alicia A. Russo
PTO Reg. No. 46,192
Attorneys for Applicant
(212) 408-2500

BAKER BOTTS LLP
30 Rockefeller Plaza
New York, NY 10112

Enclosures

MARKED UP VERSION OF TECHNICAL AMENDMENTS

Please replace paragraph on page 16, starting at line 1 and ending at line 2 with the following replacement paragraph:

end downstream the EcoRI site gaattc (SEQ ID NO:1), i.e. the lack of the 3' end from nucleotide 2014 included.

Please replace paragraph on page 16, starting at line 18 and ending at line 26 with the following replacement paragraph:

To fuse the *STM* coding region to the engrailed repressor domain the frame of the unique XbaI cloning site in pRT Ωeng had to be shifted which was achieved by inserting a GATCTCGA (SEQ ID NO:2) adapter into the upstream BamHI site, which was destroyed. The STM coding region previously amplified by reverse transcriptase PCR from Arabidopsis RNA with an 5' terminal XbaI (upstream the translation start at ATG) was inserted into the XbaI site of the adpted pRT Ωeng. The STM coding sequences used contain a BamHI site immediately preceding the natural translation stop codon, which was used subsequently to create the c-terminal GR fusion.